

# Projective Space Method for Slow Invariant Manifolds of Reactive Systems

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# Introduction

## Motivation and background

- Detailed kinetics are essential for accurate modeling of real systems.
- Reactive flow systems admit multi-scale solutions.
- Manifold methods provide a potential for computational saving.
- Slow invariant manifolds (**SIMs**) describe the asymptotic structure of reactive systems' invariant attracting trajectories.
- Current manifold construction methods either **approximate** the actual SIMs or require **a close** initial guess.

## **Long-term objective**

Create an efficient algorithm that reduces the computational cost for simulating reactive flows based on a reduction in the stiffness and dimension of the composition phase space.

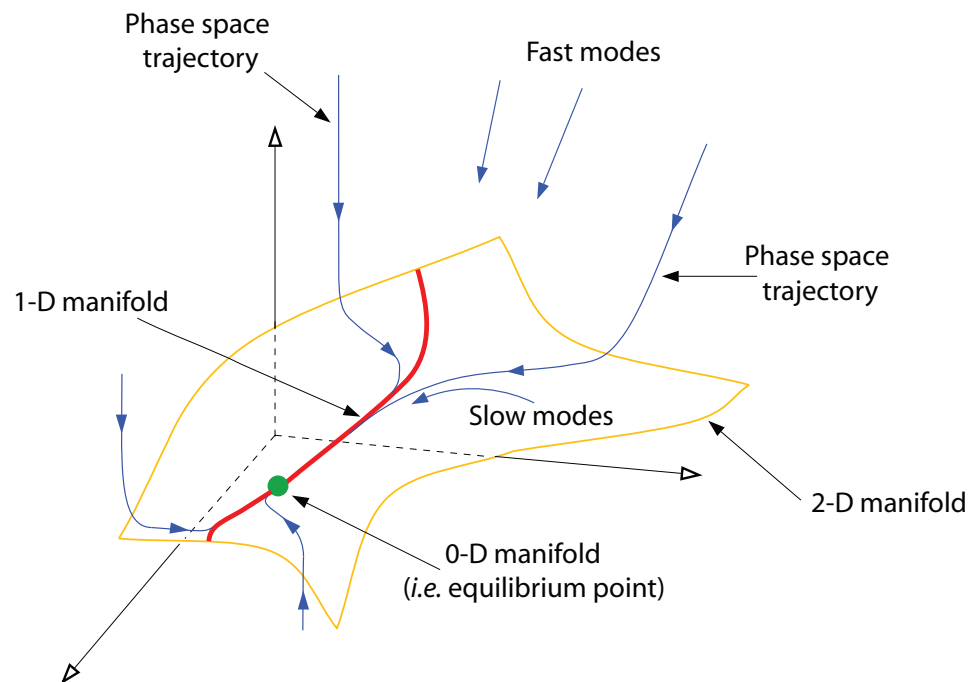
## **Immediate objective**

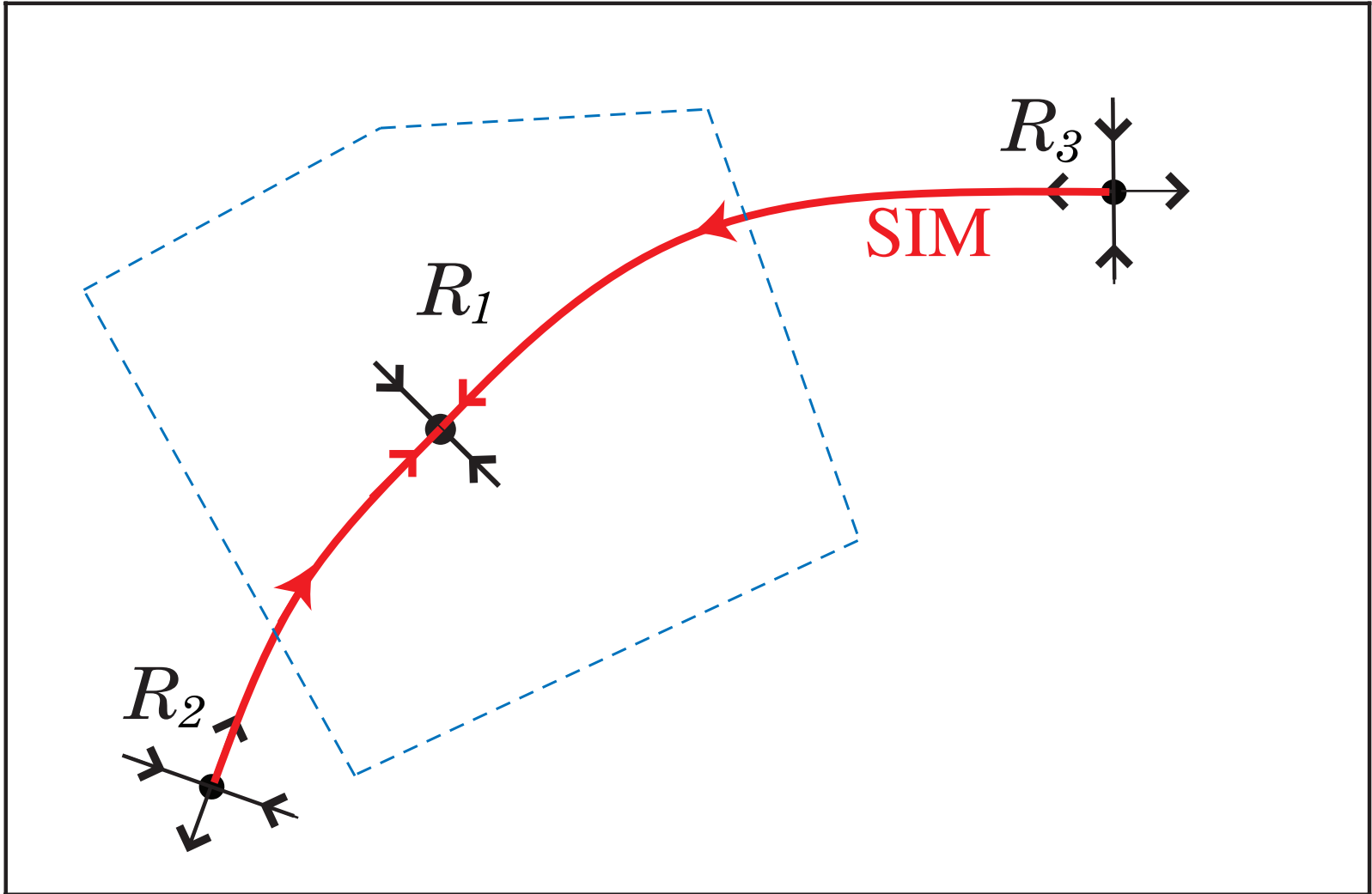
Construct 1-D SIMs for dynamical systems arising from modeling unsteady spatially homogenous closed reactive systems.

# Slow Invariant Manifold (SIM)

- The composition phase space for closed spatially homogeneous reactive system:

$$\frac{dz}{dt} = \mathbf{f}(z), \quad z \in \mathbb{R}^3.$$





# Mathematical Model

For a mixture of mass  $M$  confined in volume  $V$  containing  $N$  species composed of  $L$  elements that undergo  $J$  reversible reactions,

$$\frac{dn_i}{dt} = V \sum_{j=1}^J \nu_{ij} r_j, \quad i = 1, \dots, N,$$

where,

$$r_j = A_j T^{\beta_j} \exp\left(\frac{-E_j}{\bar{\mathcal{R}}T}\right) \left( \prod_{i=1}^N \left(\frac{n_i}{V}\right)^{\nu'_{ij}} - \frac{1}{K_j^c} \prod_{i=1}^N \left(\frac{n_i}{V}\right)^{\nu''_{ij}} \right), \quad j = 1, \dots, J,$$

$$K_j^c = \left(\frac{p^o}{\bar{\mathcal{R}}T}\right)^{\sum_{i=1}^N \nu_{ij}} \exp\left(-\frac{\sum_{i=1}^N \bar{\mu}_i^o \nu_{ij}}{\bar{\mathcal{R}}T}\right), \quad j = 1, \dots, J.$$

## System reduction

- In chemical reactions, atoms are conserved:

$$\sum_{i=1}^N \phi_{li} \nu_{ij} = 0, \quad l = 1, \dots, L, \quad j = 1, \dots, J,$$

$$\sum_{i=1}^N \phi_{li} n_i^* = \sum_{i=1}^N \phi_{li} n_i, \quad l = 1, \dots, L.$$

- Solutions of the following form exist,

$$n_i = n_i^* + M \sum_{k=1}^R \mathcal{D}_{ik} z_k, \quad i = 1, \dots, N.$$

- The reactive system is recast as an autonomous dynamical system,

$$\frac{dz_k}{dt} = f_k(z_1, \dots, z_R), \quad k = 1, \dots, R.$$

# Method of Construction

## Equilibria

- The construction method is based on identifying all the equilibria, and connecting relevant ones via heteroclinic orbits.
- For isothermal reactive systems, reaction rates depend on combinations of polynomials of  $\mathbf{z}$ .
- The set of equilibria of the full reaction network is complex:  
$$\mathbf{z}^e \in \mathbb{C}^R \mid \mathbf{f}(\mathbf{z}^e) = \mathbf{0}.$$
- This set contains **finite** and **infinite** equilibria.
- The system's equilibria can be positive dimensional continua.



## SIM construction

- A 1-D SIM has a maximum of two branches that connect two equilibria to the unique physical critical point (a sink) tangent to its slowest mode.
- These equilibria are identified by their special dynamical character: *their eigenvalue spectrum contains only one unstable direction.*
- Heteroclinic orbits are generated tangent to these special equilibria's unstable directions.
- Check first the finite equilibria, then the infinite ones.

## Projective space

- One-to-one mapping of the composition space,  $\mathbb{R}^R \rightarrow \mathbb{R}^R$ ,

$$Z_k = \frac{1}{z_k}, \quad k \in \{1, \dots, R\},$$

$$Z_i = \frac{z_i}{z_k}, \quad i \neq k, \quad i = 1, \dots, R.$$

- This maps equilibria located at infinity into a finite domain.
- To deal with the time singularity, we add the transformation

$$\frac{dt}{d\tau} = (Z_k)^{d-1},$$

where  $d$  is the highest polynomial degree of  $\mathbf{f}(\mathbf{z})$ .

## Computational strategy

- We use the `Bertini`<sup>a</sup> software (based on a homotopy continuation numerical technique) to compute the system's equilibria up to any desired accuracy.
- Thermodynamic data is obtained from `Chemkin-II`.
- The SIM heteroclinic orbits are obtained by numerical integration of the species evolution equations using a computationally inexpensive scheme.
- Computation time is typically less than 1 minute on a  $2.16\text{ GHz}$  MacBook Pro machine.

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<sup>a</sup>D. J. Bates, J. D. Hauenstein, A. J. Sommese, and C. W. Wampler, Bertini: Software for numerical algebraic geometry. Available at: [www.nd.edu/~sommese/bertini](http://www.nd.edu/~sommese/bertini).

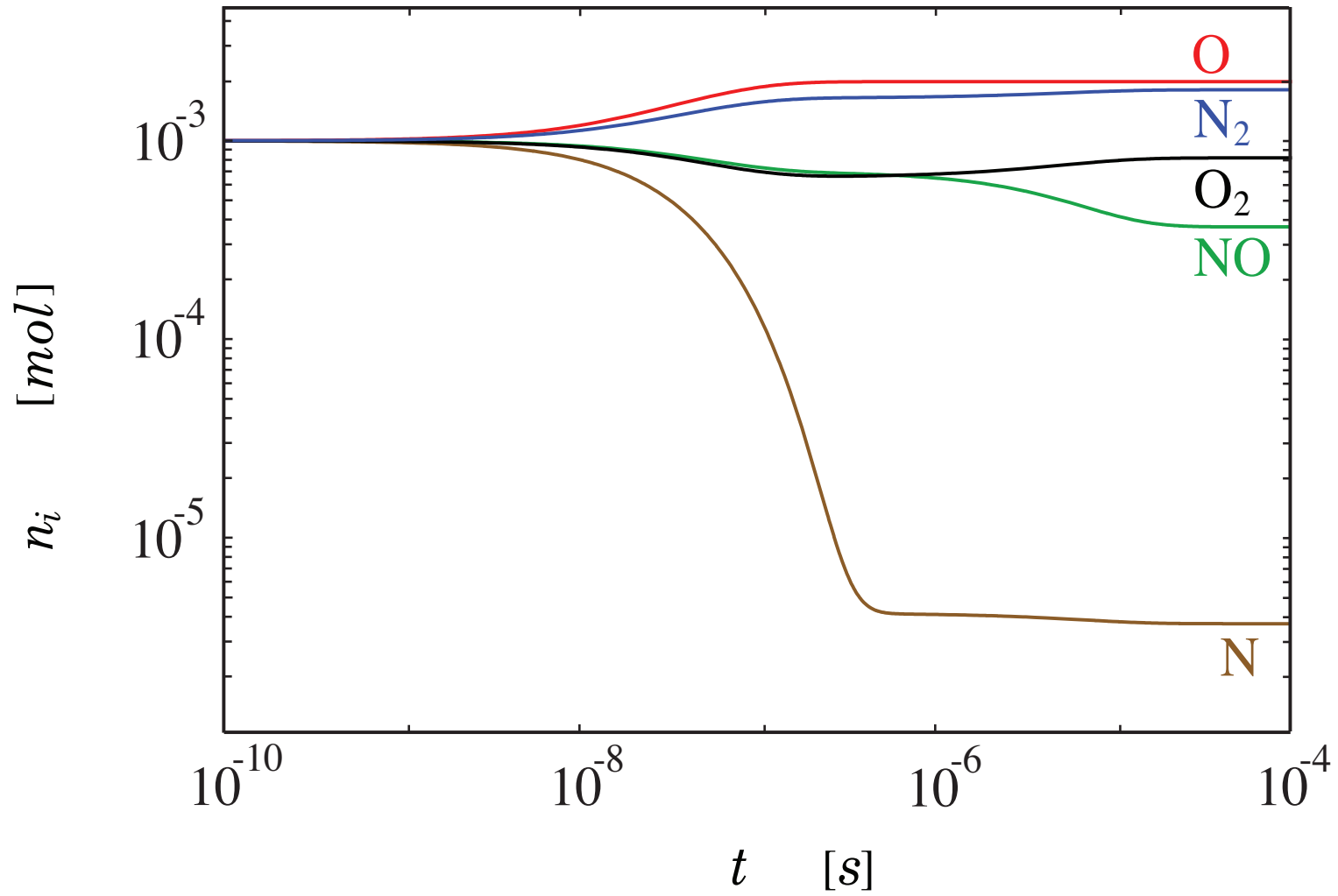
# Zel'dovich Mechanism

- The mechanism consists of  $J = 2$  bimolecular reversible reactions involving  $N = 5$  species  $\{NO, N, O, O_2, N_2\}$  and  $L = 2$  elements  $\{N, O\}$ .
- $\mathbf{z} \in \mathbb{R}^2$ , so selected species are  $i = \{1, 2\} = \{NO, N\}$ .
- The kinetic data are adopted from Baulch *et al.*<sup>b</sup>
- The system is spatially homogenous with isothermal and isochoric conditions,  $T = 4000\text{ K}$ ,  $V = 10^3\text{ cm}^3$ .
- Initial number of moles of all species are  $\mathbf{n}^* = 10^{-3}\text{ mol}$ .

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<sup>b</sup>D. L. Baulch *et al.*, *J. Phys. Chem. Ref. Data*, **34**, pp. 757-1326, 2005.

# Reactive system evolution



## Dynamical system formulation

- The evolution of the system is described by:

$$\frac{d}{dt} \begin{pmatrix} z_1 \\ z_2 \end{pmatrix} = \begin{pmatrix} 2.51 \times 10^2 + 1.16 \times 10^7 z_2 + 6.99 \times 10^8 z_2^2 \\ -9.98 \times 10^4 z_1 - 3.22 \times 10^9 z_2 z_1 \\ 2.51 \times 10^2 - 1.17 \times 10^7 z_2 - 6.98 \times 10^8 z_2^2 \\ +8.47 \times 10^4 z_1 - 1.84 \times 10^9 z_2 z_1 \end{pmatrix} \equiv \mathbf{f}(\mathbf{z}).$$

- Employ the projective space mapping with  $d = 2$  and  $k = 1$ :

$$\frac{d}{d\tau} \begin{pmatrix} t \\ Z_1 \\ Z_2 \end{pmatrix} = Z_1^2 \cdot \begin{pmatrix} Z_1^{-1} \\ -Z_1 f_1(Z_1, Z_2) \\ f_2(Z_1, Z_2) - Z_2 f_1(Z_1, Z_2) \end{pmatrix} \equiv \mathbf{F}(\mathbf{Z}),$$

## System's equilibria

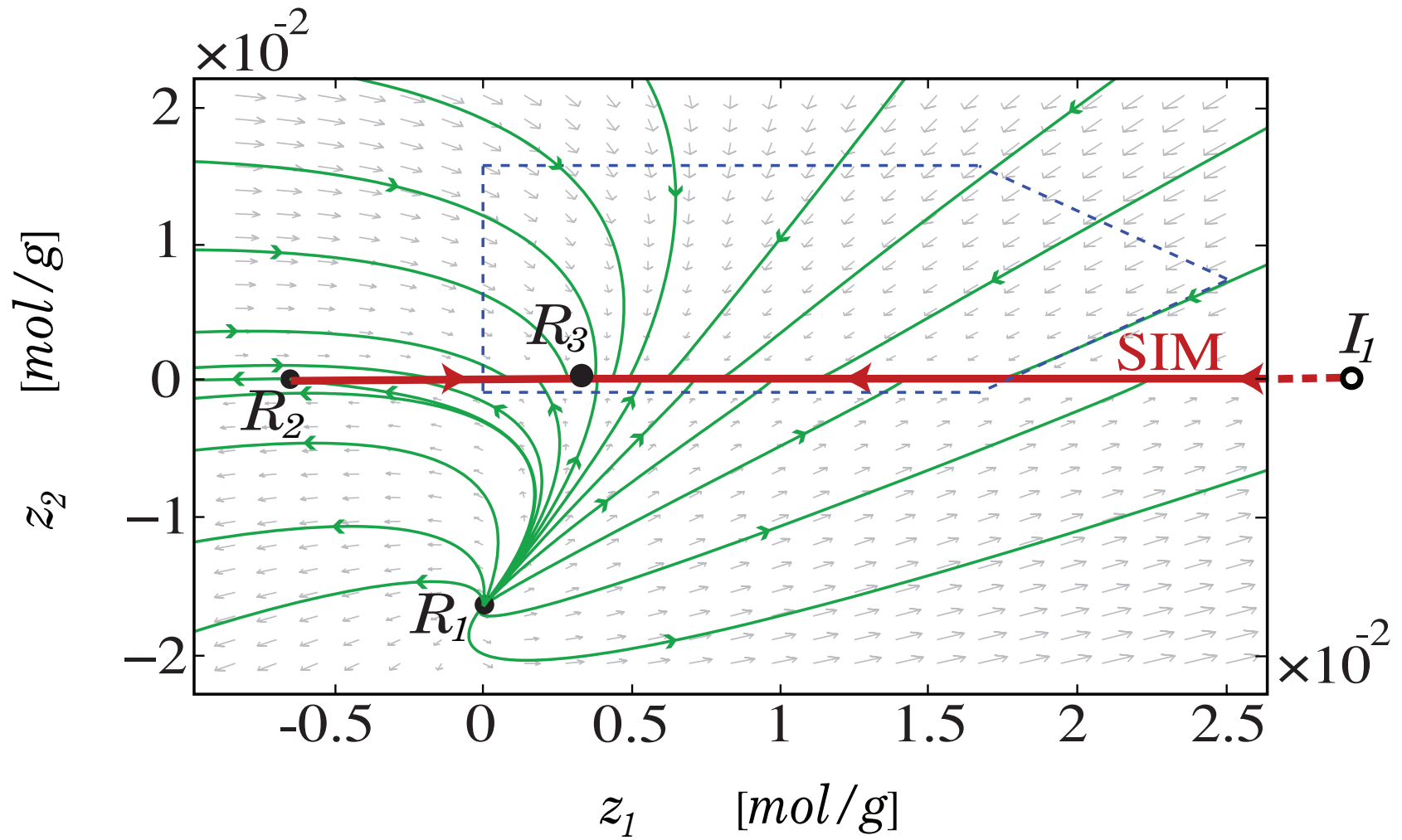
finite

$$\left\{ \begin{array}{l} R_1 \equiv (\mathbf{z}^e) = (-1.78 \times 10^{-5}, -1.67 \times 10^{-2}) \text{ mol/g}, \\ (\boldsymbol{\lambda}) = (4.18 \times 10^7, 2.35 \times 10^7) \text{ 1/s}, \\ R_2 \equiv (\mathbf{z}^e) = (-4.20 \times 10^{-3}, -2.66 \times 10^{-5}) \text{ mol/g}, \\ (\boldsymbol{\lambda}) = (-4.64 \times 10^6, 7.11 \times 10^5) \text{ 1/s}, \\ R_3 \equiv (\mathbf{z}^e) = (3.05 \times 10^{-3}, 2.94 \times 10^{-5}) \text{ mol/g}, \\ (\boldsymbol{\lambda}) = (-1.73 \times 10^7, -1.91 \times 10^5) \text{ 1/s}. \end{array} \right.$$

infinite

$$\left\{ \begin{array}{l} I_1 \equiv (\mathbf{Z}^e) = (0, 0), \\ (\boldsymbol{\lambda}) = (-1.84 \times 10^9, 0) \text{ g/mol/s}^2, \\ I_2 \equiv (\mathbf{Z}^e) = (0, 1.01), \\ (\boldsymbol{\lambda}) = (2.54 \times 10^9, 1.12 \times 10^9) \text{ g/mol/s}^2, \\ I_3 \equiv (\mathbf{Z}^e) = (0, 2.60), \\ (\boldsymbol{\lambda}) = (3.65 \times 10^9, -2.90 \times 10^9) \text{ g/mol/s}^2. \end{array} \right.$$

# The system's 1-D SIM





# Detailed Hydrogen-Air Mechanism

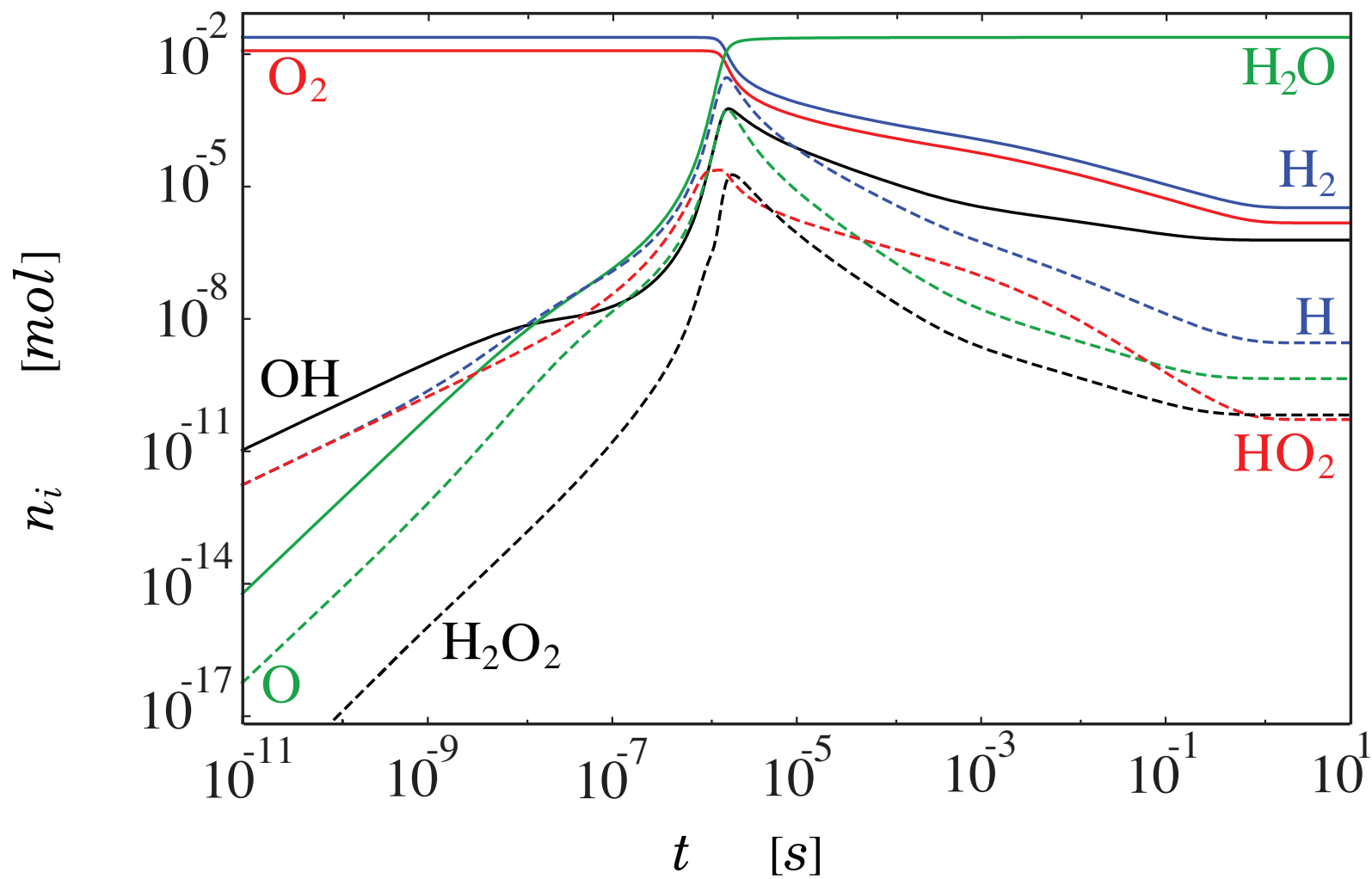
- Mechanism:  $J = 19$  reversible reactions involving  $N = 9$  species and  $L = 3$  elements.  $R = 6$ , so that  $\mathbf{z} \in \mathbb{R}^6$ .
- Kinetic model from Miller *et al.*<sup>c</sup>
- Closed and spatially homogenous system with isothermal and isochoric conditions at  $T = 1500\text{ K}$ , and  $V = 10^{-3}\text{ cm}^3$ .
- Stoichiometric mixture  $2H_2 + (O_2 + 3.76N_2)$ .
- Selected species:

$$i = \{1, 2, 3, 4, 5, 6\} = \{H_2, O_2, H, O, OH, H_2O\}.$$

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<sup>c</sup>J. A. Miller, R. E. Mitchell, M. D. Smooke, and R. J. Kee, *Proc. Combust. Ins.* **19**, pp. 181-196, 1982.

# Reactive system evolution

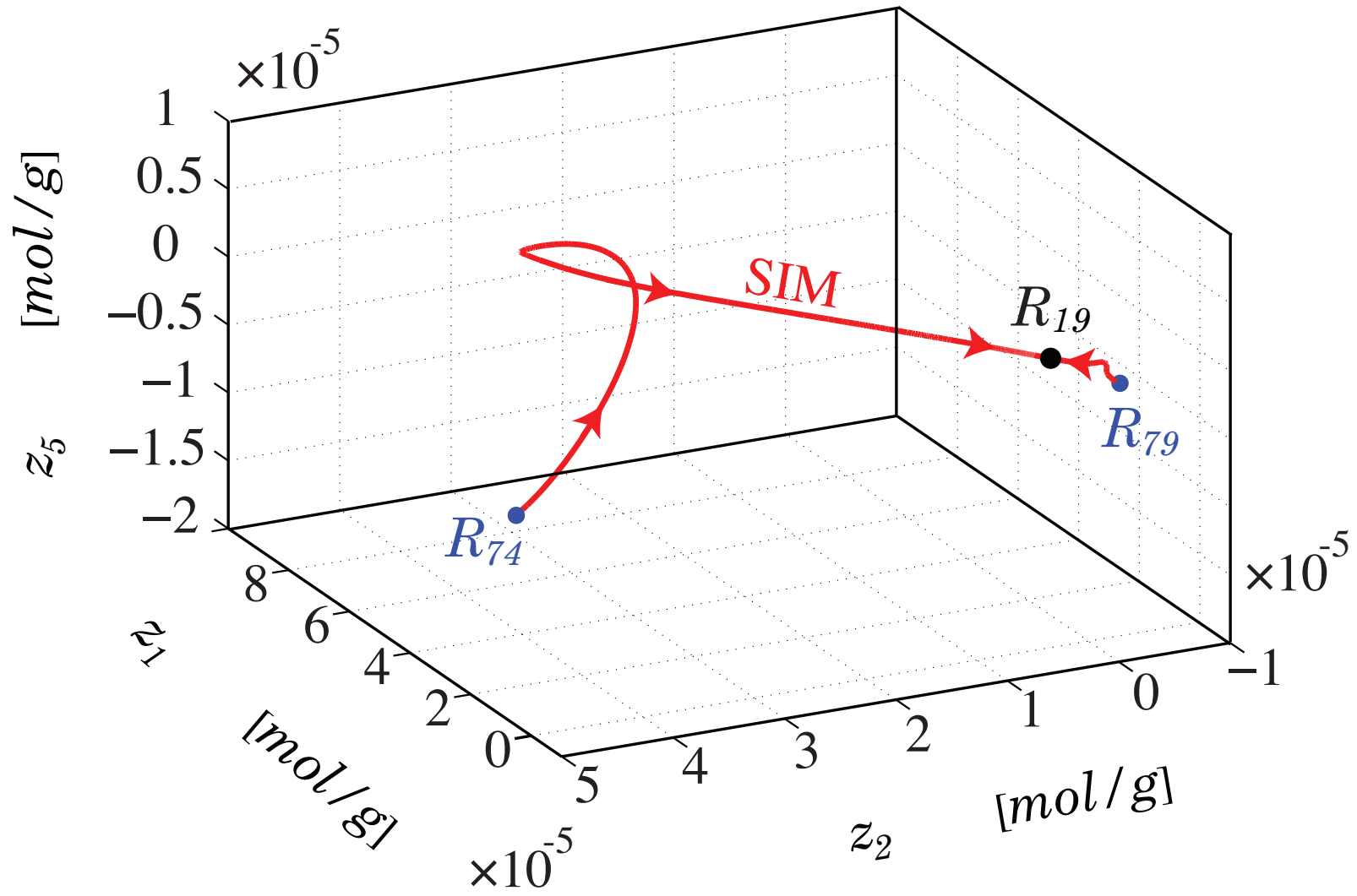


## System's equilibria

- The system has 284 **finite** and 42 **infinite** equilibria.
- The set of finite equilibria contains 90 real and 186 complex 0- $D$ , one 1- $D$ , one 2- $D$ , and six 3- $D$  equilibria.
- The set of infinite equilibria contains 18 real and 18 complex 0- $D$ , and six 1- $D$  equilibria.
- Only 14 critical points have an eigenvalue spectrum that contains only one unstable mode.
- Inside the physical domain there is a unique equilibrium:

$$R_{19} = (1.98 \times 10^{-6}, 9.00 \times 10^{-7}, 1.72 \times 10^{-9}, \\ 2.67 \times 10^{-10}, 3.66 \times 10^{-7}, 1.44 \times 10^{-2}) \text{ mol/g.}$$

### 3-D projection of the system's SIM



# Summary and Conclusions

- Once the difficult task of identifying all equilibria is complete, constructing the actual SIM is computationally **efficient** and algorithmically **easy**; thus, there is **no need** to identify it only approximately.
- Identifying all critical points, **finite** and **infinite**, plays a major role in the construction of the SIM.
- The construction procedure can be systematically extended to construct higher-dimensional SIMs.